

SCORE Search Results Details for Application 10552515 and Search Result 20080630_144103_us-10-552-515-3.ra1.

Score Home	Retrieve Application	SCORE System	SCORE	Comments /
Page	List	Overview	FAQ	Suggestions

This page gives you Search Results detail for the Application 10552515 and Search Result 20080630_144103_us-10-552-515-3.ra1.

[Go Back to previous page](#)

GenCore version 6.2.1

Copyright (c) 1993 - 2008 Bioccceleration Ltd.

OM protein - protein search, using sw model

Run on: June 30, 2008, 17:46:21 ; Search time 40 Seconds
(without alignments)
42.303 Million cell updates/sec

Title: US-10-552-515-3
Perfect score: 46
Sequence: 1 SLFMALWAV 9

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 1143754 seqs, 186252778 residues

Total number of hits satisfying chosen parameters: 1143754

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : Issued_Patents_AA:*
1: /ABSS/Data/CRF/ptodata/1/iaa/5_COMB.pep:*
2: /ABSS/Data/CRF/ptodata/1/iaa/6_COMB.pep:*
3: /ABSS/Data/CRF/ptodata/1/iaa/7_COMB.pep:*
4: /ABSS/Data/CRF/ptodata/1/iaa/H_COMB.pep:*
5: /ABSS/Data/CRF/ptodata/1/iaa/PCTUS_COMB.pep:*
6: /ABSS/Data/CRF/ptodata/1/iaa/RE_COMB.pep:*
7: /ABSS/Data/CRF/ptodata/1/iaa/backfiles1.pep:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

%

Result No.	Score	Query Match	Length	DB	ID	Description
1	40	87.0	117	3	US-10-703-032-142336	Sequence 142336,
2	39	84.8	642	3	US-10-108-260A-4483	Sequence 4483, Ap
3	36	78.3	207	2	US-08-811-519-30	Sequence 30, Appl
4	36	78.3	220	2	US-09-489-039A-13425	Sequence 13425, A
5	36	78.3	250	2	US-09-248-796A-20183	Sequence 20183, A
6	36	78.3	274	4	US-10-038-895A-1	Sequence 1, Appli
7	36	78.3	440	2	US-09-631-603-22	Sequence 22, Appl
8	36	78.3	440	2	US-09-826-509-567	Sequence 567, App
9	36	78.3	440	3	US-10-925-095-567	Sequence 567, App
10	36	78.3	442	2	US-09-538-092-637	Sequence 637, App
11	36	78.3	449	1	US-08-142-439A-5	Sequence 5, Appli
12	36	78.3	449	1	US-08-869-477-5	Sequence 5, Appli
13	35	76.1	487	2	US-09-328-352-6206	Sequence 6206, Ap
14	34	73.9	108	2	US-09-489-039A-13025	Sequence 13025, A
15	34	73.9	144	3	US-10-703-032-126625	Sequence 126625,
16	34	73.9	152	2	US-09-489-039A-11538	Sequence 11538, A
17	34	73.9	218	2	US-09-270-767-42075	Sequence 42075, A
18	34	73.9	435	2	US-09-252-991A-19124	Sequence 19124, A
19	34	73.9	968	3	US-09-252-691C-7784	Sequence 7784, Ap
20	33	71.7	169	2	US-10-094-749-1824	Sequence 1824, Ap
21	33	71.7	202	3	US-10-703-032-125681	Sequence 125681,
22	33	71.7	225	3	US-09-540-209B-7498	Sequence 7498, Ap
23	33	71.7	240	3	US-10-703-032-136346	Sequence 136346,
24	33	71.7	252	3	US-10-369-493-7925	Sequence 7925, Ap
25	33	71.7	362	3	US-10-369-493-4227	Sequence 4227, Ap
26	33	71.7	469	2	US-09-328-352-5007	Sequence 5007, Ap
27	33	71.7	507	3	US-10-369-493-10701	Sequence 10701, A
28	33	71.7	524	2	US-09-252-991A-18580	Sequence 18580, A
29	33	71.7	528	3	US-09-602-740-34	Sequence 34, Appl
30	33	71.7	530	3	US-09-602-740-32	Sequence 32, Appl
31	33	71.7	575	3	US-10-805-394A-4263	Sequence 4263, Ap
32	33	71.7	595	3	US-10-703-032-120251	Sequence 120251,
33	33	71.7	596	2	US-10-104-047-2541	Sequence 2541, Ap
34	33	71.7	920	2	US-10-104-047-2574	Sequence 2574, Ap
35	33	71.7	1280	3	US-10-343-657-7	Sequence 7, Appli
36	33	71.7	1359	3	US-10-736-769-44	Sequence 44, Appl
37	32	69.6	86	3	US-10-198-232-64	Sequence 64, Appl
38	32	69.6	126	3	US-10-198-232-61	Sequence 61, Appl
39	32	69.6	157	3	US-10-703-032-137302	Sequence 137302,
40	32	69.6	169	3	US-10-703-032-210817	Sequence 210817,
41	32	69.6	226	2	US-09-248-796A-20804	Sequence 20804, A
42	32	69.6	256	3	US-10-038-854-177	Sequence 177, App
43	32	69.6	270	2	US-09-134-000C-5024	Sequence 5024, Ap
44	32	69.6	291	2	US-09-902-540-15052	Sequence 15052, A
45	32	69.6	298	2	US-09-602-777A-384	Sequence 384, App

ALIGNMENTS

RESULT 1

US-10-703-032-142336

; Sequence 142336, Application US/10703032

```

; Patent No. 7214786
; GENERAL INFORMATION:
; APPLICANT: Kovalic, David K.
; APPLICANT: Andersen, Scott E.
; APPLICANT: Byrum, Joseph R.
; APPLICANT: Conner, Timothy W.
; APPLICANT: Cao, Yongwei
; APPLICANT: Masucci, James D.
; APPLICANT: Zhou, Yihua
; TITLE OF INVENTION: Nucleic Acid Molecules And Other Molecules Associated With
; TITLE OF INVENTION: Plants
; FILE REFERENCE: 38-21(53374)B
; CURRENT APPLICATION NUMBER: US/10/703,032
; CURRENT FILING DATE: 2003-11-06
; PRIOR APPLICATION NUMBER: 10/020,338
; PRIOR FILING DATE: 2001-12-12
; NUMBER OF SEQ ID NOS: 211164
; SEQ ID NO 142336
; LENGTH: 117
; TYPE: PRT
; ORGANISM: Triticum aestivum
; FEATURE:
; OTHER INFORMATION: Clone ID: PAT_TA_36754.pep
US-10-703-032-142336

```

```

Query Match          87.0%; Score 40; DB 3; Length 117;
Best Local Similarity 77.8%; Pred. No. 19;
Matches      7; Conservative      2; Mismatches      0; Indels      0; Gaps      0;

```

```

Qy      1 SLFMALWAV 9
        |:|:|||||
Db      96 SIFIALWAV 104

```

RESULT 2

```

US-10-108-260A-4483
; Sequence 4483, Application US/10108260A
; Patent No. 7193069
; GENERAL INFORMATION:
; APPLICANT: HELIX RESEARCH INSTITUTE
; TITLE OF INVENTION: No. 7193069e1 full length cDNA
; FILE REFERENCE: H1-A0106
; CURRENT APPLICATION NUMBER: US/10/108,260A
; CURRENT FILING DATE: 2002-03-27
; NUMBER OF SEQ ID NOS: 5458
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 4483
; LENGTH: 642
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-108-260A-4483

```

```

Query Match          84.8%; Score 39; DB 3; Length 642;
Best Local Similarity 87.5%; Pred. No. 1.5e+02;
Matches      7; Conservative      1; Mismatches      0; Indels      0; Gaps      0;

```

Qy 1 SLFMALWA 8
 |:|||||
 Db 385 SVFMALWA 392

RESULT 3

US-08-811-519-30
 ; Sequence 30, Application US/08811519B
 ; Patent No. 6630345
 ; GENERAL INFORMATION:
 ; APPLICANT: Petrenko, Alexandre
 ; TITLE OF INVENTION: CALCIUM INDEPENDENT RECEPTOR OF ALPHA-LATROTOXIN,
 ; TITLE OF INVENTION: CHARACTERIZATION AND USES THEREOF
 ; FILE REFERENCE: 1049-1-007
 ; CURRENT APPLICATION NUMBER: US/08/811,519B
 ; CURRENT FILING DATE: 1997-03-04
 ; NUMBER OF SEQ ID NOS: 31
 ; SOFTWARE: PatentIn Ver. 2.0
 ; SEQ ID NO 30
 ; LENGTH: 207
 ; TYPE: PRT
 ; ORGANISM: rat
 US-08-811-519-30

Query Match 78.3%; Score 36; DB 2; Length 207;
 Best Local Similarity 55.6%; Pred. No. 1.5e+02;
 Matches 5; Conservative 4; Mismatches 0; Indels 0; Gaps 0;

Qy 1 SLFMALWAV 9
 ::|:||||:
 Db 110 AIFVALWAI 118

RESULT 4

US-09-489-039A-13425
 ; Sequence 13425, Application US/09489039A
 ; Patent No. 6610836
 ; GENERAL INFORMATION:
 ; APPLICANT: Gary Breton et. al
 ; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO KLEBSIELLA
 ; TITLE OF INVENTION: PNEUMONIAE FOR DIAGNOSTICS AND THERAPEUTICS
 ; FILE REFERENCE: 2709.2004001
 ; CURRENT APPLICATION NUMBER: US/09/489,039A
 ; CURRENT FILING DATE: 2000-01-27
 ; PRIOR APPLICATION NUMBER: US 60/117,747
 ; PRIOR FILING DATE: 1999-01-29
 ; NUMBER OF SEQ ID NOS: 14342
 ; SEQ ID NO 13425
 ; LENGTH: 220
 ; TYPE: PRT
 ; ORGANISM: Klebsiella pneumoniae
 US-09-489-039A-13425

Query Match 78.3%; Score 36; DB 2; Length 220;
 Best Local Similarity 77.8%; Pred. No. 1.6e+02;

Matches 7; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

Qy 1 SLFMALWAV 9
 |||| || |
 Db 9 SLFMKLWLV 17

RESULT 5

US-09-248-796A-20183

; Sequence 20183, Application US/09248796A

; Patent No. 6747137

; GENERAL INFORMATION:

; APPLICANT: Keith Weinstock et al

; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO CANDIDA ALBICANS

; TITLE OF INVENTION: FOR DIAGNOSTICS AND THERAPEUTICS

; FILE REFERENCE: 107196.132

; CURRENT APPLICATION NUMBER: US/09/248,796A

; CURRENT FILING DATE: 1999-02-12

; PRIOR APPLICATION NUMBER: US 60/074,725

; PRIOR FILING DATE: 1998-02-13

; PRIOR APPLICATION NUMBER: US 60/096,409

; PRIOR FILING DATE: 1998-08-13

; NUMBER OF SEQ ID NOS: 28208

; SEQ ID NO 20183

; LENGTH: 250

; TYPE: PRT

; ORGANISM: Candida albicans

US-09-248-796A-20183

Query Match 78.3%; Score 36; DB 2; Length 250;

Best Local Similarity 77.8%; Pred. No. 1.8e+02;

Matches 7; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

Qy 1 SLFMALWAV 9
 || :|||||
 Db 70 SLIIALWAV 78

RESULT 6

US-10-038-895A-1

; Sequence 1, Application US/10038895A

; Patent No. H002136

; GENERAL INFORMATION:

; APPLICANT: Kulp, David C.

; APPLICANT: Siani-Rose, Michael A.

; APPLICANT: Williams, Alan J.

; APPLICANT: Harmon, Cyrus L.

; TITLE OF INVENTION: Nucleic Acids Encoding G Proteins Coupled Receptors

; FILE REFERENCE: 3379.1

; CURRENT APPLICATION NUMBER: US/10/038,895A

; CURRENT FILING DATE: 2003-03-25

; PRIOR APPLICATION NUMBER: US 60/244,082

; PRIOR FILING DATE: 2000-10-26

; NUMBER OF SEQ ID NOS: 20

; SOFTWARE: PatentIn version 3.2

```

; SEQ ID NO 1
; LENGTH: 274
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Synthetic Organism
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION: (126)..(126)
; OTHER INFORMATION: Xaa can be any naturally occurring amino acid
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION: (146)..(146)
; OTHER INFORMATION: Xaa can be any naturally occurring amino acid
US-10-038-895A-1

```

```

Query Match          78.3%; Score 36; DB 4; Length 274;
Best Local Similarity 55.6%; Pred. No. 2e+02;
Matches      5; Conservative      4; Mismatches      0; Indels      0; Gaps      0;

```

```

Qy      1 SLFMALWAV 9
        ::|:||||:
Db      170 AIFVALWAI 178

```

RESULT 7

US-09-631-603-22

```

; Sequence 22, Application US/09631603
; Patent No. 6733990
; GENERAL INFORMATION:
; APPLICANT: Hodge, Martin R.
; APPLICANT: Lloyd, Clare
; APPLICANT: Weich, Nadine
; TITLE OF INVENTION: 15571, A No. 6733990e1 GPCR-like Molecule of the
; TITLE OF INVENTION: Secretin-Like Family and Uses Thereof
; FILE REFERENCE: 5800-48A
; CURRENT APPLICATION NUMBER: US/09/631,603
; CURRENT FILING DATE: 2000-08-03
; PRIOR APPLICATION NUMBER: 09/515,781
; PRIOR FILING DATE: 2000-02-29
; PRIOR APPLICATION NUMBER: 60/146,916
; PRIOR FILING DATE: 2000-08-03
; NUMBER OF SEQ ID NOS: 24
; SOFTWARE: FastSEQ for Windows Version 3.0
; SEQ ID NO 22
; LENGTH: 440
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-631-603-22

```

```

Query Match          78.3%; Score 36; DB 2; Length 440;
Best Local Similarity 55.6%; Pred. No. 3.2e+02;
Matches      5; Conservative      4; Mismatches      0; Indels      0; Gaps      0;

```

```

Qy      1 SLFMALWAV 9
        ::|:||||:

```

Db 267 AIFVALWAI 275

RESULT 8

US-09-826-509-567

; Sequence 567, Application US/09826509

; Patent No. 6806054

; GENERAL INFORMATION:

; APPLICANT: Lehmann-Bruinsma, Karin

; APPLICANT: Liaw, Chen W.

; APPLICANT: Lin, I-Lin

; TITLE OF INVENTION: No. 6806054-Endogenous, Constitutively Activated Known G

; TITLE OF INVENTION: Protein-Coupled Receptors

; FILE REFERENCE: AREN-207

; CURRENT APPLICATION NUMBER: US/09/826,509

; CURRENT FILING DATE: 2001-04-05

; PRIOR APPLICATION NUMBER: 60/195,747

; PRIOR FILING DATE: 2000-04-07

; PRIOR APPLICATION NUMBER: 09/170,496

; PRIOR FILING DATE: 1998-10-13

; NUMBER OF SEQ ID NOS: 589

; SOFTWARE: PatentIn Version 2.1

; SEQ ID NO 567

; LENGTH: 440

; TYPE: PRT

; ORGANISM: Homo sapiens

US-09-826-509-567

Query Match 78.3%; Score 36; DB 2; Length 440;

Best Local Similarity 55.6%; Pred. No. 3.2e+02;

Matches 5; Conservative 4; Mismatches 0; Indels 0; Gaps 0;

Qy 1 SLFMALWAV 9

::|:||||:

Db 267 AIFVALWAI 275

RESULT 9

US-10-925-095-567

; Sequence 567, Application US/10925095

; Patent No. 7097969

; GENERAL INFORMATION:

; APPLICANT: Lehmann-Bruinsma, Karin

; APPLICANT: Liaw, Chen W.

; APPLICANT: Lin, I-Lin

; TITLE OF INVENTION: No. 7097969-Endogenous, Constitutively Activated Known G

; TITLE OF INVENTION: Protein-Coupled Receptors

; FILE REFERENCE: AREN-207

; CURRENT APPLICATION NUMBER: US/10/925,095

; CURRENT FILING DATE: 2004-08-24

; PRIOR APPLICATION NUMBER: US/09/826,509

; PRIOR FILING DATE: 2001-04-05

; PRIOR APPLICATION NUMBER: 60/195,747

; PRIOR FILING DATE: 2000-04-07

; PRIOR APPLICATION NUMBER: 09/170,496

```
; PRIOR FILING DATE: 1998-10-13
; NUMBER OF SEQ ID NOS: 589
; SOFTWARE: PatentIn Version 2.1
; SEQ ID NO 567
; LENGTH: 440
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-925-095-567
```

```
Query Match          78.3%; Score 36; DB 3; Length 440;
Best Local Similarity 55.6%; Pred. No. 3.2e+02;
Matches      5; Conservative      4; Mismatches      0; Indels      0; Gaps      0;
```

```
Qy      1 SLFMALWAV 9
        ::|:||||:
Db      267 AIFVALWAI 275
```

RESULT 10

US-09-538-092-637

```
; Sequence 637, Application US/09538092
; Patent No. 6753314
; GENERAL INFORMATION:
; APPLICANT: Giot, Loic
; APPLICANT: Mansfield, Traci A.
; TITLE OF INVENTION: Protein-Protein Complexes and Method of Using Same
; FILE REFERENCE: 15966-542
; CURRENT APPLICATION NUMBER: US/09/538,092
; CURRENT FILING DATE: 2000-03-29
; PRIOR APPLICATION NUMBER: 60/127,352
; PRIOR FILING DATE: 1999-04-01
; PRIOR APPLICATION NUMBER: 60/178,965
; PRIOR FILING DATE: 2000-02-01
; NUMBER OF SEQ ID NOS: 1387
; SOFTWARE: CuraPatSeqFormatter Version 0.9
; SEQ ID NO 637
; LENGTH: 442
; TYPE: PRT
; ORGANISM: Saccharomyces cerevisiae
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION: (0)...(0)
; OTHER INFORMATION: Polypeptide Accession Number YMR243C
US-09-538-092-637
```

```
Query Match          78.3%; Score 36; DB 2; Length 442;
Best Local Similarity 77.8%; Pred. No. 3.2e+02;
Matches      7; Conservative      1; Mismatches      1; Indels      0; Gaps      0;
```

```
Qy      1 SLFMALWAV 9
        || :|||||
Db      48 SLLVALWAV 56
```

RESULT 11

US-08-142-439A-5


```

; Sequence 5, Application US/08142439A
; Patent No. 5670360
; GENERAL INFORMATION:
; APPLICANT: Thorens, Bernard
; TITLE OF INVENTION: Receptor for the Glucagon-Like-Peptide-1
; TITLE OF INVENTION: (GLP-1)
; NUMBER OF SEQUENCES: 9
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: No. 5670360o No. 5670360disk of No. 5670360th America, Inc.
; STREET: 405 Lexington Avenue, Suite 6400
; CITY: New York
; STATE: New York
; COUNTRY: U.S.A.
; ZIP: 10174-6201
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/142,439A
; FILING DATE: 24-NOV-93
; CLASSIFICATION: 530
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: DK 398/92
; FILING DATE: 25-MAR-92
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: PCT/EP93/00697
; FILING DATE: 23-MAR-93
; ATTORNEY/AGENT INFORMATION:
; NAME: Harrington, James J.
; REGISTRATION NUMBER: 38,711
; REFERENCE/DOCKET NUMBER: 3756.204-US
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 212 867 0123
; TELEFAX: 212 867 0298
; INFORMATION FOR SEQ ID NO: 5:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 449 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: protein
; HYPOTHETICAL: NO
; ANTI-SENSE: NO
; ORIGINAL SOURCE:
; ORGANISM: Rattus norvegicus
; STRAIN: Sprague-Dawley
US-08-142-439A-5

```

Query Match 78.3%; Score 36; DB 1; Length 449;
 Best Local Similarity 55.6%; Pred. No. 3.3e+02;
 Matches 5; Conservative 4; Mismatches 0; Indels 0; Gaps 0;

Qy 1 SLFMALWAV 9

::|:||||:

Db

267 AIFVALWAI 275

RESULT 12

US-08-869-477-5

; Sequence 5, Application US/08869477

; Patent No. 5846747

; GENERAL INFORMATION:

; APPLICANT: Thorens, Bernard

; TITLE OF INVENTION: Receptor for the Glucagon-Like-Peptide-1

; TITLE OF INVENTION: (GLP-1)

; NUMBER OF SEQUENCES: 9

; CORRESPONDENCE ADDRESS:

; ADDRESSEE: No. 5846747o No. 5846747disk of No. 5846747th America, Inc.

; STREET: 405 Lexington Avenue, Suite 6400

; CITY: New York

; STATE: New York

; COUNTRY: U.S.A.

; ZIP: 10174-6201

; COMPUTER READABLE FORM:

; MEDIUM TYPE: Floppy disk

; COMPUTER: IBM PC compatible

; OPERATING SYSTEM: PC-DOS/MS-DOS

; SOFTWARE: PatentIn Release #1.0, Version #1.25

; CURRENT APPLICATION DATA:

; APPLICATION NUMBER: US/08/869,477

; FILING DATE:

; CLASSIFICATION: 435

; PRIOR APPLICATION DATA:

; APPLICATION NUMBER: US/08/142,439

; FILING DATE: 24-NOV-93

; APPLICATION NUMBER: DK 398/92

; FILING DATE: 25-MAR-92

; PRIOR APPLICATION DATA:

; APPLICATION NUMBER: PCT/EP93/00697

; FILING DATE: 23-MAR-93

; ATTORNEY/AGENT INFORMATION:

; NAME: Harrington, James J.

; REGISTRATION NUMBER: 38,711

; REFERENCE/DOCKET NUMBER: 3756.204-US

; TELECOMMUNICATION INFORMATION:

; TELEPHONE: 212 867 0123

; TELEFAX: 212 867 0298

; INFORMATION FOR SEQ ID NO: 5:

; SEQUENCE CHARACTERISTICS:

; LENGTH: 449 amino acids

; TYPE: amino acid

; STRANDEDNESS: single

; TOPOLOGY: linear

; MOLECULE TYPE: protein

; HYPOTHETICAL: NO

; ANTI-SENSE: NO

; ORIGINAL SOURCE:

; ORGANISM: Rattus norvegicus

; STRAIN: Sprague-Dawley

US-08-869-477-5

Query Match 78.3%; Score 36; DB 1; Length 449;
 Best Local Similarity 55.6%; Pred. No. 3.3e+02;
 Matches 5; Conservative 4; Mismatches 0; Indels 0; Gaps 0;

Qy 1 SLFMALWAV 9
 ::l:||||:
 Db 267 AIFVALWAI 275

RESULT 13

US-09-328-352-6206

; Sequence 6206, Application US/09328352

; Patent No. 6562958

; GENERAL INFORMATION:

; APPLICANT: Gary L. Breton et al.

; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO ACINETOBACTER

; TITLE OF INVENTION: BAUMANNII FOR DIAGNOSTICS AND THERAPEUTICS

; FILE REFERENCE: GTC99-03PA

; CURRENT APPLICATION NUMBER: US/09/328,352

; CURRENT FILING DATE: 1999-06-04

; NUMBER OF SEQ ID NOS: 8252

; SEQ ID NO 6206

; LENGTH: 487

; TYPE: PRT

; ORGANISM: Acinetobacter baumannii

US-09-328-352-6206

Query Match 76.1%; Score 35; DB 2; Length 487;
 Best Local Similarity 85.7%; Pred. No. 5.2e+02;
 Matches 6; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

Qy 1 SLFMALW 7
 ||||:||
 Db 54 SLFMSLW 60

RESULT 14

US-09-489-039A-13025

; Sequence 13025, Application US/09489039A

; Patent No. 6610836

; GENERAL INFORMATION:

; APPLICANT: Gary Breton et. al

; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO KLEBSIELLA

; TITLE OF INVENTION: PNEUMONIAE FOR DIAGNOSTICS AND THERAPEUTICS

; FILE REFERENCE: 2709.2004001

; CURRENT APPLICATION NUMBER: US/09/489,039A

; CURRENT FILING DATE: 2000-01-27

; PRIOR APPLICATION NUMBER: US 60/117,747

; PRIOR FILING DATE: 1999-01-29

; NUMBER OF SEQ ID NOS: 14342

; SEQ ID NO 13025

; LENGTH: 108

; TYPE: PRT

; ORGANISM: *Klebsiella pneumoniae*
 US-09-489-039A-13025

Query Match 73.9%; Score 34; DB 2; Length 108;
 Best Local Similarity 75.0%; Pred. No. 1.7e+02;
 Matches 6; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

Qy 1 SLFMALWA 8

|||: |||

Db 86 SLFLRLWA 93

RESULT 15

US-10-703-032-126625

; Sequence 126625, Application US/10703032

; Patent No. 7214786

GENERAL INFORMATION:

; APPLICANT: Kovalic, David K.

; APPLICANT: Andersen, Scott E.

; APPLICANT: Byrum, Joseph R.

; APPLICANT: Conner, Timothy W.

; APPLICANT: Cao, Yongwei

; APPLICANT: Masucci, James D.

; APPLICANT: Zhou, Yihua

; TITLE OF INVENTION: Nucleic Acid Molecules And Other Molecules Associated With

; TITLE OF INVENTION: Plants

; FILE REFERENCE: 38-21(53374)B

; CURRENT APPLICATION NUMBER: US/10/703,032

; CURRENT FILING DATE: 2003-11-06

; PRIOR APPLICATION NUMBER: 10/020,338

; PRIOR FILING DATE: 2001-12-12

; NUMBER OF SEQ ID NOS: 211164

; SEQ ID NO 126625

; LENGTH: 144

; TYPE: PRT

; ORGANISM: *Triticum aestivum*

; FEATURE:

; OTHER INFORMATION: Clone ID: PAT_TA_21043.pep

US-10-703-032-126625

Query Match 73.9%; Score 34; DB 3; Length 144;
 Best Local Similarity 66.7%; Pred. No. 2.3e+02;
 Matches 6; Conservative 1; Mismatches 2; Indels 0; Gaps 0;

Qy 1 SLFMALWAV 9

|:| ||||

Db 51 SVFFCLWAV 59

Search completed: June 30, 2008, 17:51:38

Job time : 39.625 secs